Figure 1

SEQ ID NO:1

- 1 MKLVSVALMY LGSLAFLGAD TARLDVASEF RKKWNKWALS RGKRELRMSS SYPTGLADVK
- 61 AGPAQTLIRP ODMKGASRSP EDSSPDAARI RVKRYRQSMN NFQGLRSFGC RFGTCTVQKL
- 121 AHQIYQFTDK DKDNVAPRSK ISPQGYGRRR RRSLPEAGPG RTLVSSKPQA HGAPAPPSGS
- 181 APHFL

SEQ ID NO:2

- 1 ARLDVASEFR KKWNKWALSR GKRELRMSSS YPTGLADVKA GPAQTLIRPQ DMKGASRSPE
- 61 DSSPDAARIR VKRYRQSMNN FQGLRSFGCR FGTCTVQKLA HQIYQFTDKD KDNVAPRSKI
- 121 SPQGYGRRRR RSLPEAGPGR TLVSSKPQAH GAPAPPSGSA PHFL

SEQ ID NO:3

ELRMSSSYPT GLADVKAGPA QTLIRPQDMK GASRSPEDSS PDAARIRV

SEQ ID NO:4

MSSSYPTGLA DVKAGPAQTL IRPQDMK

SEQ ID NO:5

MSSSYPTGLA DVK

SEQ ID NO:6

AGPAQTLIRP QDMK